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=> s 16 (3A) (cell line)
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          3641 L6 (3A) (CELL LINE)
=> s 17 and 14
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             1 L7 AND L4 - test hydrox / P450/ C4P 3A4
                            human Diver/Lepatocyke / cell line
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     124:75473
     An investigation of the interaction between halofantrine, CYP2D6 and
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     expression systems
     Halliday, Rachel C.; Jones, Barry C.; Smith, Dennis A.; Kitteringham, Neil
ΑU
     R.; Park, B. Kevin
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     British Journal of Clinical Pharmacology (1995), 40(4), 369-78
SO
     CODEN: BCPHBM; ISSN: 0306-5251
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     English
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     The authors have assessed the interaction of the antimalarial halofantrine
     with cytochrome P 450 (CYP) enzymes in vitro, with the use of microsomes
     from human liver and recombinant cell
     lines. Rac-halofantrine was a potent inhibitor (IC50 = 1.06
     .mu.M, Ki \approx 4.3 .mu.M) of the 1-hydroxylation of bufuralol, a marker for
     CYP2D6 activity. Of a group of structurally related antimalarials tested,
     only quinidine (IC50 = 0.04 .mu.M) was more potent. Microsomes prepd.
     from recombinant CYP2D6 and CYP3A4 cell lines were shown to catalyze
     halofantrine N-debutylation. The metab. of halofantrine to its N-desbutyl
     metabolite by human liver microsomes showed no correlation with CYP2D6
     genotypic or phenotypic status and there was no consistent inhibition by
     quinidine. The rate of halofantrine metab. showed a significant
     correlation with both CYP3A4 protein levels (r = 0.88) and the rate of
     felodipine metab. (r = 0.86), a marker substrate for CYP3A4 activity.
     Inhibition studies showed that ketoconazole is a potent inhibitor of
     halofantrine metab. (IC50 = 1.57 \cdot mu.M). In conclusion, the authors have
     demonstrated that halofantrine is a potent inhibitor of CYP2D6 in vitro
     and can also be metabolized by the enzyme. However, in human liver
     microsomes it appears to be metabolized largely by CYP3A4.
<---->User Break---->
<---->
=> s transfection (3a) (17 or hepg2)
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